

WHAT IS CLAIMED IS:

- 1                   1.       A method of detecting cancer cells in a biological sample from a  
2 mammal, the method comprising steps of:  
3                   (i) providing the biological sample from the mammal; and  
4                   (ii) detecting a nucleic acid molecule encoding a PRC17 polypeptide  
5 comprising at least 85% amino acid sequence identity to an amino acid sequence of SEQ  
6 ID NO:2 or at least 70% amino acid identity to an amino acid sequence of SEQ ID NO:4  
7 or SEQ ID NO:6 in the biological sample, wherein an increase in the level of the nucleic  
8 acid molecule in the sample compared to normal indicates the presence of cancer cells.
- 1                   2.       The method of claim 1, wherein the polypeptide has an amino acid  
2 sequence of SEQ ID NO:2, SEQ ID NO:4 or SEQ ID NO:6.
- 1                   3.       The method of claim 1, wherein the detecting step further  
2 comprises:  
3                   (a) contacting the nucleic acid molecule with a probe under conditions in  
4 which the probe selectively hybridizes to the nucleic acid molecule to form a stable  
5 hybridization complex; and  
6                   (b) detecting the hybridization complex.
- 1                   4.       The method of claim 3, wherein the contacting step further  
2 comprises a step of amplifying the gene in an amplification reaction.
- 1                   5.       The method of claim 4, wherein the amplification reaction is a  
2 polymerase chain reaction.
- 1                   6.       The method of claim 1, wherein the nucleic acid is an mRNA.
- 1                   7.       The method of claim 1, wherein the biological sample is a tissue  
2 biopsy.
- 1                   8.       The method of claim 7, wherein the cancer cells are selected from  
2 the group consisting of prostate tissue, breast tissue, lung tissue, and ovarian tissue.
- 1                   9.       The method of claim 1, wherein the mammal is a human.

1 10. A method of detecting a presence of cancer cells in a biological  
2 sample from a mammal, the method comprising steps of:

3 (i) providing the biological sample from the mammal; and

4 (ii) detecting an overexpression of a polypeptide comprising polypeptide  
5 comprising at least 85% amino acid sequence identity to an amino acid sequence of SEQ  
6 ID NO:2 or at least 70% amino acid identity to an amino acid sequence of SEQ ID NO:4  
7 or SEQ ID NO:6 in the biological sample, thereby detecting the presence of cancer cells  
8 in the biological sample.

1 11. The method of claim 10, wherein the polypeptide has an amino  
2 acid sequence of SEQ ID NO:2, SEQ ID NO:4 or SEQ ID NO:6.

1 12. The method of claim 10, wherein the polypeptide is detected using an  
2 antibody that selectively binds to the polypeptide.

1 13. The method of claim 10, wherein the biological sample is a tissue  
2 biopsy.

1 14. The method of claim 10, wherein the cancer cells are selected from  
2 the group consisting of prostate cancer cells, breast cancer cells, lung cancer cells, and  
3 ovarian cancer cells.

1 15. The method of claim 10, wherein the mammal is a human.

1 16. A method of monitoring the efficacy of a therapeutic treatment of a  
2 cancer, the method comprising the steps of:

3 (i) providing a biological sample from a mammal undergoing the  
4 therapeutic treatment; and

5 (ii) detecting a level of a polypeptide comprising at least 85% amino acid  
6 sequence identity to an amino acid sequence of SEQ ID NO:2 or at least 70% amino acid  
7 identity to an amino acid sequence of SEQ ID NO:4 or SEQ ID NO:6 in the biological  
8 sample compared to a level in a biological sample from the mammal prior to, or earlier in,  
9 the therapeutic treatment, thereby monitoring the efficacy of the therapy.

1 17. The method of claim 16, wherein the polypeptide has an amino  
2 acid sequence of SEQ ID NO:2, SEQ ID NO:4 or SEQ ID NO:6.

1 18. The method of claim 16, wherein the cancer is selected from the  
2 group consisting of prostate cancer, ovarian cancer, lung cancer, and breast cancer.

1 19. The method of claim 16, wherein the polypeptide is detected using  
2 an antibody that selectively binds to the polypeptide.

1 20. A method of monitoring the efficacy of a therapeutic treatment of a  
2 cancer, the method comprising the steps of:

3 (i) providing a biological sample from a mammal undergoing the  
4 therapeutic treatment; and

5 (ii) detecting a nucleic acid molecule encoding a PRC17 polypeptide  
6 comprising at least 85% amino acid sequence identity to an amino acid sequence of SEQ  
7 ID NO:2 or at least 70% amino acid identity to an amino acid sequence of SEQ ID NO:4  
8 or SEQ ID NO:6 in the biological sample compared to a level in a biological sample from  
9 the mammal prior to, or earlier in, the therapeutic treatment, thereby monitoring the  
10 efficacy of the therapy.

1 21. An isolated nucleic acid encoding a PRC17 polypeptide, the  
2 nucleic acid encoding a polypeptide comprising at least 85% amino acid identity to an  
3 amino acid sequence of SEQ ID NO:2 or at least 70% identity to an amino acid sequence  
4 of SEQ ID NO:4 or SEQ ID NO:6.

1 22. The nucleic acid of claim 21, wherein the nucleic acid encodes a  
2 PRC17 polypeptide that specifically binds to polyclonal antibodies generated against an  
3 amino acid sequence of SEQ ID NO:2, SEQ ID NO:4 or SEQ ID NO:6.

1 23. The nucleic acid of claim 21, wherein the nucleic acid encodes a  
2 PRC17 polypeptide comprising an amino acid sequence of SEQ ID NO:2, SEQ ID NO:4  
3 or SEQ ID NO:6.

1 24. The nucleic acid of claim 23, wherein the nucleic acid comprises a  
2 nucleotide sequence of SEQ ID NO:1, SEQ ID NO:3 or SEQ ID NO:5.

1 25. The nucleic acid of claim 21, wherein the nucleic acid is amplified  
2 by primers that specifically hybridize under stringent hybridization conditions to a nucleic  
3 acid having a nucleotide sequence of SEQ ID NO:1, SEQ ID NO:3 or SEQ ID NO:5.

1                   26.     The nucleic acid of claim 21, wherein the nucleic acid specifically  
2     hybridizes under stringent hybridization conditions to a nucleic acid having a nucleotide  
3     sequence of SEQ ID NO:1, SEQ ID NO:3 or SEQ ID NO:5.

1                   27.     An isolated PRC17 polypeptide, the polypeptide comprising at  
2     least 85% amino acid sequence identity to an amino acid sequence of SEQ ID NO:2 or at  
3     least 70% amino acid identity to an amino acid sequence of SEQ ID NO:4 or SEQ ID  
4     NO:6.

1                   28.     The isolated polypeptide of claim 8, wherein the polypeptide  
2     specifically binds to polyclonal antibodies generated against SEQ ID NO:2, SEQ ID  
3     NO:4 or SEQ ID NO:6.

1                   29.     The isolated polypeptide of claim 8, wherein the polypeptide has  
2     an amino acid sequence of SEQ ID NO:2, SEQ ID NO:4 or SEQ ID NO:6.

1                   30.     An antibody that selectively binds to the polypeptide of claim 8.

1                   31.     An expression vector comprising the nucleic acid of claim 1.

1                   32.     A host cell transfected with the vector of claim 31.

1                   33.     A method of identifying a compound that modulates activity of a  
2     PRC17 polypeptide, the method comprising steps of:

3                   (i) contacting the polypeptide with the compound, wherein the polypeptide  
4     comprises at least 85% amino acid sequence identity to an amino acid sequence of SEQ  
5     ID NO:2 or at least 70% amino acid identity to an amino acid sequence of SEQ ID NO:4  
6     or SEQ ID NO:6; and

7                   (ii) determining the functional effect of the compound on the polypeptide.

1                   34.     The method of claim 33, wherein the polypeptide is linked to a  
2     solid phase.

1                   35.     The method of claim 33, wherein the polypeptide is expressed in a  
2     cell or cell membrane.

10071553-5555  
200909-5555

1                   36.     The method of claim 33, wherein the polypeptide has an amino  
2     acid sequence of SEQ ID NO:2, SEQ ID NO:4 or SEQ ID NO:6.

1                   37.     A method of treating a disease or condition associated with the  
2     activity of a PRC17 polypeptide, the method comprising the step of administering to a  
3     subject a therapeutically effective amount of a compound identified using the method of  
4     claim 33.

1                   38.     The method of claim 37, wherein the subject is a human.

1                   39.     The method of claim 18, wherein the compound is an antibody.

1                   40.     A method of inhibiting proliferation of a cancer cell that expresses a  
2     polypeptide comprising at least 85% amino acid sequence identity to an amino acid sequence  
3     of SEQ ID NO:2 or at least 70% amino acid identity to an amino acid sequence of SEQ ID  
4     NO:4 or SEQ ID NO:6, the method comprising the step of contacting the cancer cell with a  
5     therapeutically effective amount of an inhibitor of the polypeptide.

1                   41.     The method of claim 40, wherein the polypeptide has an amino acid  
2     sequence of SEQ ID NO:2, SEQ ID NO:4 or SEQ ID NO:6.

1                   42.     The method of claim 40, wherein the cancer cell is selected from  
2     the group consisting of a prostate cancer cell, a breast cancer cancer cell, a lung cancer  
3     cell or an ovarian cancer cell.

1                   43.     The method of claim 40, wherein the inhibitor is an antibody.